

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/572,882
Source: IFWP
Date Processed by STIC: 3/31/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 03/31/2006

PATENT APPLICATION: US/10/572,882

TIME: 12:19:12

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\03292006\J572882.raw

3 <110> APPLICANT: Bayer HealthCare AG
 4 Golz, Stefan
 5 Broggemeier, Ulf
 6 Geerts, Andreas
 8 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases
 Associated with
 9 G-Protein Coupled Receptor ADIPOR2 (ADIPOR2)
 11 <130> FILE REFERENCE: LeA.36 902
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/572,882
 C--> 13 <141> CURRENT FILING DATE: 2006-03-20
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/010383
 14 <151> PRIOR FILING DATE: 2004-09-16
 16 <150> PRIOR APPLICATION NUMBER: EP03021898.6
 17 <151> PRIOR FILING DATE: 2003-09-27
 19 <160> NUMBER OF SEQ ID NOS: 5
 21 <170> SOFTWARE: PatentIn version 3.3
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 3500
 25 <212> TYPE: DNA
 26 <213> ORGANISM: Homo sapiens
 28 <400> SEQUENCE: 1
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 47 tcagtatcgg ggagtaagag caggagtgtt tttgggccta ggctgagtg gaatcattcc 600
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 63 gggctctgcg tggtagatga ctgagaagag aaaaacaaaa ataaatcata cctcaaagga 1080
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 67 ctactgattg cgggctctgc aagacccttg gcaaaactggc ttctgatcca tatcatattt 1200
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 73 tgattttaaa tgctcttttg ggagaacaaa gaaattaatg taaataagat ttctaactgt 1380

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143 gttccattga tgtgggattt ccctacttgc tgtattctca gtttctaata aaaagaacca 3480
145 aatgaaaaaa aaaaaaaaaa 3500
148 <210> SEQ ID NO: 2
149 <211> LENGTH: 258
150 <212> TYPE: PRT
151 <213> ORGANISM: Homo sapiens
153 <400> SEQUENCE: 2
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156 1 5 10 15
159 Glu Thr Gly Asn Ile Trp Thr His Leu Leu Gly Cys Val Phe Phe Leu
160 20 25 30
163 Cys Leu Gly Ile Phe Tyr Met Phe Arg Pro Asn Ile Ser Phe Val Ala
164 35 40 45
167 Pro Leu Gln Glu Lys Val Val Phe Gly Leu Phe Phe Leu Gly Ala Ile
168 50 55 60

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171 Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr Val Tyr Cys His Ser
172 65                               70                               75                               80
175 Glu Gly Val Ser Arg Leu Phe Ser Lys Leu Asp Tyr Ser Gly Ile Ala
176                               85                               90                               95
179 Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu Tyr Tyr Ser Phe Tyr
180                               100                              105                              110
183 Cys Asn Pro Gln Pro Cys Phe Ile Tyr Leu Ile Val Ile Cys Val Leu
184                               115                              120                              125
187 Gly Ile Ala Ala Ile Ile Val Ser Gln Trp Asp Met Phe Ala Thr Pro
188                               130                              135                              140
191 Gln Tyr Arg Gly Val Arg Ala Gly Val Phe Leu Gly Leu Gly Leu Ser
192 145                               150                              155                              160
195 Gly Ile Ile Pro Thr Leu His Tyr Val Ile Ser Glu Gly Phe Leu Lys
196                               165                              170                              175
199 Ala Ala Thr Ile Gly Gln Ile Gly Trp Leu Met Leu Met Ala Ser Leu
200                               180                              185                              190
203 Tyr Ile Thr Gly Ala Ala Leu Tyr Ala Ala Arg Ile Pro Glu Arg Phe
204                               195                              200                              205
207 Phe Pro Gly Lys Cys Asp Ile Trp Phe His Ser His Gln Leu Phe His
208                               210                              215                              220
211 Ile Phe Val Val Ala Gly Ala Phe Val His Phe His Gly Val Ser Asn
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223 <210> SEQ ID NO: 3

224 <211> LENGTH: 20

225 <212> TYPE: DNA

226 <213> ORGANISM: Artificial

228 <220> FEATURE:

229 <223> OTHER INFORMATION: Primer1 (forward primer)

231 <400> SEQUENCE: 3

232 catggtgtct caaacctcca

20

235 <210> SEQ ID NO: 4

236 <211> LENGTH: 20

237 <212> TYPE: DNA

238 <213> ORGANISM: Artificial

240 <220> FEATURE:

241 <223> OTHER INFORMATION: Primer2 (reverse primer)

243 <400> SEQUENCE: 4

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252 <220> FEATURE:

253 <223> OTHER INFORMATION: Probe1

255 <400> SEQUENCE: 5

256 agtttcgttt catgatcggc ggg

23

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/572,882

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Input Set : A:\PTO.KD.txt
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#: 3, 4, 5

VERIFICATION SUMMARY

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Input Set : A:\PTO.KD.txt

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date